Sequence Listing

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SEO ID
                rat OGFr cDNA
          1
    5
          2
                rat OGFr protein
                rat OGFr cDNA partial (clone 14)
          3
          4
                human OGFr cDNA (Provisional)
                human OGFr splice version 8, cDNA
          5
          6
                human OGFr splice version 8, protein
    10
                human OGEr splice version 1, cDNA
          7
          8
                human OGF splice version 1, protein
               human OGFr\splice version 4, cDNA
          9
                human OGFr splice version 4, protein
          10
          11
                human OGFr splice version 7, cDNA
                human OGFr splice version 7, protein
    15
          12
               human OGFr splace version 127, cDNA
          13
                human OGFr splide version 127, protein
          14
rat OGFr antisense
                                          5'-GACTCAGGGACTTAGCTTCATCC-3'
          15
                                          5'-ATAGATACTACGCCGGCTGTCCT-3'
          16
                scrambled
                                          5'-GGTCGTCCATGCTCGGCTAGAAT-3'
    20
          17
                human OGFr antisense
                scrambled
                                          5'-GTGCAGTGCAATGCTCTCCGTGA-3'
          18
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- Rat Opioid Growth Factor Receptor cDNA sequence SEO ID NO: 1

TGGGCTCAGCCACGCCCCAGGGTGCCCCCAGTGGGACTAGTTCTTCATTCTGGCAGCTGCACACATCTGTCAGTGAGGGAATGTCAGGTC 90 TCTCACTCTCCTCTCACTATCCTTTCCGCAGAAAGCGGGTCCTCCTGCTTGTCGAGTATGGACGACCCGGACTGCGATTCCACCTGG 180 GAGGAGGAGGAGGAGGAGGAGGATGGCCAGGCGGATGATACGACCGATGAGGACACGGGCGACGATGACGCGGACGCGGAGGAG 270 GCACGGCCAAGCCTGTTCCAGTCCAGGATGACAGGGTACCGAAACTGGCGTGCTATGCAGGACATGCAAAGATACCGGCACAACTACCCG 360 GATTTGACAGATCAAGACTGdAATGGTGACATGTGCAACCTGAGCTTCTACAAAAATGAGATCTGCTTCCAGCCAAATGGGGCTCTCATC 450 GAGGACATTCTTCAGAACTGGAAGAACTATGACCTCCTGGAAGAGAATCACTCCTACATCCAGTGGCTGTTTCCTCTGCGGGAACCA 540 GGAGTGAACTGGCACGCCAAGCCCTCACCCTGAAGGAGGTTGAGGCATTTAAAAGCTCCAAGGAAGTCAGAGAGCGTCTTGTCCGGGCC 630 TATGAGCTCATGCTGGGCTTCTATGGGTTCCACCTTGAGGACCGGGGCACGGGTGCTGTATGCCGTGCACAGAACTTCCAGCCGCGCTTC 720 CACAATCTGAACAGCCACAGCCACAACCATGCGTATTACACGCATCCTCAAGTCACTGGGTGAGCTGGGCTTAGAACACTACCAGGCACCCCTGGTCCGCTTCTTCCTGGAGGAGACCCTTGTACAGCACAAACTGCCCAGCGTGCGCCAGAGTGCCCTGGACTACTTCCTGTTCGCT 900 GTGCGCTGCCGGCACCAGCGCCGGGAGCTTGTGTACTTTGCCTGGGAGCACTTCAAGCCTCGCCGAGAGTTTGTCTGGGGGCCCCGTGAC990 AAGCTGCGGAGATTCAAGCCCCAGAQCATACCCCAGCCACTGACGGGACCAGGGCAGGCAGATAAAGATGAGGGCTCCAGGGACCCCTCC 1080 CAAGAGGCTGGCACCCAGGGTCGGAC¢TGTGGATCTGGAAGGGACCTGAGTGGGGACAGTGGAACAGCTGAGGATCCCTCACTGCTGAAC 1170 ACAAAGCCCTCAGATGGGGAACCTTGGATGGGAACCAGAGGGATGAAGCTAAGTCCCTGAGTCCCAAGGAGGAGAAAAAGGAAGTTG 1260 GAGGGGAACAGGCAGGAGCAGGTCCCAdGGGAGGCAGATCCCCAGGGTGTCTCTGAGGTAGAGAAAATTGCCCTTAACCTTGAGGAGTGT 1350 GCCCTTAGCCCTATCAGCCAGGAGCCCAGGAGGCTGAACCGCCCTGTCCTGTGGCCAGGGTGGCTAATGAGGTAAGAAAGCGGAGGAAG 1440 GTGGAGGAAGGGGCTGAGGGTGATGGAGTAGTAACACTCAAATGCAGGCCAGTGCCCTGCCTACCCCTTCAGAGTGTCCTGAG 1530 GCCCAAAAGGATGGGAATGGGCCAGAGGACQCAAACAGCCAGGTTGGGGCAGAGGATTCCAAAAGCCAGGTGGGGCCGGAGGATCCAAAC 1620 AGCCAGGTGGGGCTGGAGGACCCAAACAGCQAGGTCGGGCCAGAGGACCCAAACAGCCAGGTCGGGCCAGAGGACCCAAACAGCCAGGTC 1710 GGGCCAGAGGACCCAAACAGCCAGGTCGGGCQAGAGAGCCCAAACAGCCAGGTGGTGGGGGCCAGAGCAAGCTGCCTCTAAGAGCCCTGTG 1800 GAGGACCCTGACTCTGACACTATGGGAACCTCAGTGGATGAGTCAGAGGAGTTGGCAAGGATTGAGGCCTCTGCTGAACCCCCAAAGCCT 1890 TAGAGGTGCATCTCAGTCCTACTCAGCCCACTGCAGGGGGTTTCTGAGTCCAGAGCTCTGCCGTAGGCTCTTCTTGGTGCCCCACAGTGC 1980 $\tt TGGCCTCTCCCTAGTGGTCACTGAGGTGGCCAC\ref{thm:constraint} Agastraint and the transfer of the tr$

Initiator AUG at 151-153 Terminator TAG at 1891-1983 Open reading frame 151 (AUGGACGAC...) to 1890 (... AAGCCT)

SEQ ID NO:2 -- Rat Opioid Growth Factor Receptor

- MDDPDCDSTW EEESEEDGED \GQADDTTDED TGDDDGDAEE ARPSLFQSRM TGYRNWRAMQ DMQRYRHNYP NLTDQDCNGD MCNLSFYKNE ICFQPNGALI 101 EDILONWKDN YDLLEENHSY IOWLFPLREP GVNWHAKPLT LKEVEAFKSS 151 KEVRERLVRA YELMLGFYGF HLEDRGTGAV CRAQNFQPRF HNLNSHSHNN
- 40 LRITRILKSL GELGLEHYQA PLVRFFLEET LVQHKLPSVR QSALDYFLFA

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	251	VRCRHQRREL VYFAWEHFKP RREFVWGPRD KLRRFKPQTI PQPLTGPGQA
	301	DKDEGSRDPS QEAGTQGRTC GSGRDLSGDS GTAEDPSLLN TKPSDGGTLD
	351	GNQRDEAKSL SPKESKKRKL EGNRQEQVPG EADPQGVSEV EKIALNLEEC
	401	ALSPISQEPR EAEPPCPVAR VANEVRKRRK VEEGAEGDGV VSNTQMQASA
5	451	LPPTPSECRE AQKDGNGPED SNSQVGAEDS KSQVGPEDPN SQVGLEDPNS
	501	QVGPEDPNSQ VGPEDPNSQV GPEDPNSQVG PEDPNSQVVG PEQAASKSPV
	551	EDPDSDTMGT\SVDESEELAR IEASAEPPKP
10	SEQ ID NO	:3 Rat OGFn, partial cDNA sequence, clone 14
	1	CATTGGGCCG ACGTCGCATG CTCCTCTAGA CTCGAGGAAT TCGGGCCCCA
	51	GGGTGTCTCT GAGGTAQAGA AAATTGCCCT TAACCTTGAG GAGTGTGCCC
	101	TTAGCCCTAT CAGCCAGGAG CCCAGGGAGG STGAACCGCC CTGTCCTGTG
15	151	GCCAGGGTGG CTANAATGAG GTAAGAAANG CGGNAGGAAG GTGGAGGAAG

GGGCTGAGGG TGNATGGAGT AGTCAGTAAC ACTYAAATGN CAGGCCAGTG 201 251 CCCTGCCTCC TACCCCTTCA GAGTGTCCTG AGGCCCAAAA GGATGGGAAT GGGCCAGAGG ACTCAAACAG\CCAGGTTGGG GCAGAGGATT CCAAAAGCCA 301 GGTGGGGCCG GAGGATCCAA ACAGCCAGGT GGGGCTGGAG GACCCAAACA 351 GCCAGGTCGG GCCAGAGGAC CCAAACAGCC AGGTCGGGCC AGAGGACCCA 401 AACAGCCAGG TCGGGCCAGA GGACCCAAAC AGCCAGGTCG GGCCAGAGGA 451 CCCAAACAGC CAGGTGGTGG GGQCAGAGCA AGCTGCCTCT AAGAGCCCTG 501 TGGANGGACC CTGACTCTGA CACTATGGGA ACCTCAGTGG ATGAGTCAGA 551 GGAGTTGGCA AGGATTGAGG CNTYTGCTGA ACCCCCAAAG CCTTAGAGGT 601 GCATTTCAGT CCTACTCAGC CCACTGCAGG GGGTTTCTGA GTCCAGAGCT 651 CTGCCGTAGG CTCTTCTTGG TGCCCCACAG TGCTGGCCTC TCCCTASTGG 701 TCACTGAGGT GGCCACCAGA GGGACTQAGG CCCTGCCCTC AGGGAAGGCC 751 AAGGCCTTCA GAACCCTCCT TACCTCAQTG TGTCCTCCTC CACTGCCCTC 801 TGAGCCCTGC GTTGTGATCA GACCCTAAGG GTCTAGAGGG AGGGGCCTCT 851 TCATTAGTCT GGTGCCAAGT GAGGCCTTTT CTGAATAAAC TCTTTAGACT 901 TTGTCAAAAA AAAAAAAAA AAAAAAAAAAAAAAAA 951

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SEQ ID NO: 4 -- Human OGF receptor cDNA; spliced form A Length: 2290

1 TAGAATTCAG CGGCCGCTGA ATTCTAGCCG AGCATGGACG ACCCCGACTG 51 CGACTCCAC¢ TGGGAGGAGG ACGAGGAGGA TGCGGAGGAC GCGGAGGACG 5 101 AGGACTGCGA\GGACGGCGAG GCCGCCGGCG CGAGGGACGC GGACGCAGGG 151 GACGAGGACG AGGAGTCGGA GGAGCCGCGG GCGGCGCGCG CCAGCTCGTT 201 CCAGTCCAGA ATGACAGGGT CCAGAAACTG GCGAGCCACG AGGGACATGT 251 GTAGGTATCG GQACAACTAT CCGGATCTGG TGGAACGAGA CTGCAATGGG 301 GACACGCCAA ACQTGAGTTT CTACAGAAAT GAGATCCGCT TCCTGCCCAA 10 351 CGGCTGTTTC ATTGAGGACA TTCTTCAGAA CTGGACGGAC AACTATGACC 401 TCCTTGAGGA CAATCACTCC TACATCCAGT GGCTGTTTCC TCTGCGAGAA 451 CCAGGAGTGA ACTGGCATGC CAAGCCCCTC ACGCTCAGGG AGGTCGAGGT 501 GTTTAAAAGC TCCCAGGAGA TCCAGGAGCG GCTTGTCCGG GCCTACGAGC 15 551 TCATGCTGGG CTTCTAGGGG ATCCGGCTGG AGGACCGAGG CACGGGCACG 601 GTGGGCCGAG CACAGAAQTA CCAGAAGCGC TTCCAGAACC TGAACTGGCG 651 CAGCCACAAC AACCTCCG&A TCACACGCAT CCTCAAGTCG CCGTGTGAGC 701 TGAGCCTCGA GCACTTCCAG GCGCCACTGG TCCGCTTCTT CCTGGAGGAG 751 ACGCTGGTGC GGCGGGAGCT GCCGGGGGTG CGGCAGAGTG CCCTGGACTA 20 801 CTTCATGTTC GCCGTGCGCT GCCGACACCA GCGCCGCCAG CTGGTGCACT 851 TCGCCTGGGA GCACTTCCGG &CCCGCTGCA AGTTCGTCTG GGGGCCCCAA 901 GACAAGCTGC GGAGGTTCAA GCCCAGCTCT CTGCCCCATC CGCTCGAGGG 951 CTCCAGGAAG GTGGAGGAGG AAGGAAGCCC CGGGGACCCC GACCACGAGG 1001 CCAGCACCCA GGGTCGGACC TGTGGGCCAG AGCATAGCAA GGGTGGGGGC 1051 AGGGTGGACG AGGGGCCCCA GCCACGGAGC GTGGAGCCCC AGGATGCGGG 1101 ACCCCTGGAG AGGAGCCAGG GGGATGAGGC AGGGGGCCAC GGGGAAGATA 1151 GGCCGGAGCC CTTAAGCCCC AAAGAGAGCA AGAAGAGGAA GCTGGAGCTG 1201 AGCCGGCGG AGCAGCCGCC CACAGAGCCA GGCCCTCAGA GTGCCTCAGA 1251 GGTGGAGAAG ATCGCTCTGA ATTTGGAGGG GTGTGCCCTC AGCCAGGGCA 1301 GCCTCAGGAC GGGGACCCAG GAAGTGGGCG GTCAGGACCC TGGGGAGGCA 30 1351 GTGCAGCCCT GCCGCCAACC CCTGGGAQCC AGGGTGGCCG ACAAGGTGAG 1401 GAAGCGGAGG AAGGTGGATG AGGGTGCT GGGACAGTGCT GCGGTGGCCA 1451 GTGGTGGTGC CCAGACCTTG GCCCTTGCCG GGTCCCCTGC CCCATCGGGG 1501 CACCCCAAGG CTGGACACAG TGAGAACGGG\GTTGAGGAGG ACACAGAAGG 1551 TCGAACGGGG CCCAAAGAAG GTACCCCTGG GAGCCCCATCG GAGACCCCAG 35

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1601 GCCCCAGCC AGCAGGACCT GCAGGGGACG AGCCAGCCGA GAGCCCATCG
1651 GAGACCCCAG GCCCCGCCC GGCAGGACCT GCAGGGGACG AGCCAGCCGA
1701 GAGCCCATCG GAGACCCCAG GCCCCAGCCC GGCAGGACCT ACAAGGGATG
1751 AGCCAGCCCA GAGCCCATCG GAGACCCCAG GCCCCGCCC GGCAGGACCT
5 1801 GCAGGGGACC AGCCAGCCG GAGACCCCAG GCCCCCGCCC
1851 GGCAGGACCT GCAGGGGACG AGCCAGCCG GAGACCCCAG GCCCCCGCC
1851 GCCCCAGCCC GGCAGGACCT ACAAGGGATG AGCCAGCCAA GGCGGGGGAG
1901 GCCCCAGCCC GGCAGGACCT ACAAGGGATG AGCCAGCCAA GGCGGGGGAG
1951 GCAGCAGAGT TGCAGKACGC AGAGGTGAG TCTTCTGCCA AGTCTGGGAA
2001 GCCTTAAGGA AAGGAGTGCC CGTCGGCGTC TTGGTCCTCC TGTCCCTGCT
10 2051 GCAGGGGCTG GGCCTCCGG AGCTTGCTGC GGGCTCCCCT CAGGCTCTGC
2101 TTCGTGACCC GTCACCCATG ACCCACAGTG CTGGCCTCCT GTGGGCCAC
2151 TATAGCARSC ACCAGAAGCC GCGAGGCCCT CAGGGAAGCC CAAGGCCTGC
2201 AGAAGCCTCC TGGCCTGGCT GTGTCTTCCC CACCCAGCTC TCCCCTGCGC
2251 CCCTGTCTTT GTAAATTGAC CCTTCTGGAG TGGGGGGCGG
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Letter"S", "R", or "K" was used in positions where there was a 50-50 split on the consensus sequences. S = C or G; R = A or G; K = T or G. Initiator AUG 34-36 Terminator TAA 2005-2007

20 Open reading frame: 34 \(AUGGACGAC...\) to 2004 (...GGGAAGCCT)

SEQ ID NO: 5 -- Human Opioid Growth Factor Receptor cDNA, spliced version 8

1 TAGAATTCAG CGGCCGGTGA ATTCTAGCCG AGCATGGACG ACCCCGACTG
51 CGACTCCACC TGGGAGGAGG ACGAGGAGGA TGCGGAGGAC GCGGAGGACG
101 AGGACTGCGA GGACGGCGAG GCCGCCGGCG CGAGGGACGC GGACGCAGGG
151 GACGAGGACG AGGAGTCGGA GGAGCCGCGG GCGGCGCGC CCAGCTCGTT
201 CCAGTCCAGA ATGACAGGGT CCAGAAACTG GCGAGCCACG AGGGACATGT
251 GTAGGTATCG GCACAACTAT CCGGATCTGG TGGAACGAGA CTGCAATGGG
301 GACACGCCAA ACCTGAGTTT CTACAGAAAT GAGATCCGCT TCCTGCCCAA
351 CGGCTGTTC ATTGAGGACA TTCTTCAGAA CTGGACGGAC AACTATGACC

		1				
	401	TCCTTGAGGA	CAATCACTCC	TACATCCAGT	GGCTGTTTCC	TCTGCGAGAA
	451	CCAGGAGTGA	ACTGGCATGC	CAAGCCCCTC	ACGCTCAGGG	AGGTCGAGGT
	501	GTTTAAAAGC	TCCCAGGAGA	TCCAGGAGCG	GCTTGTCCGG	GCCTACGAGC
	551	TCATGCTGGG	CTTCTACGGG	ATCCGGCTGG	AGGACCGAGG	CACGGGCACG
5	601	GTGGGCCGAG	CACAGAACTA	CCAGAAGCGC	TTCCAGAACC	TGAACTGGCG
	651	CAGCCACAAC	AACCTCCGCA	TCACACGCAT	CCTCAAGTCG	CCGTGTGAGC
	701	TGAGCCTCGA	GCACTTCCAG	GCGCCACTGG	TCCGCTTCTT	CCTGGAGGAG
	751	ACGCTGGTGC	GGCGGGAGCT	GCCGGGGGTG	CGGCAGAGTG	CCCTGGACTA
	801	CTTCATGTTC	CCGTGCGCT	GCCGACACCA	GCGCCGCCAG	CTGGTGCACT
10	851	TCGCCTGGGA	GCACTTCCGG	CCCCGCTGCA	AGTTCGTCTG	GGGGCCCCAA
	901	GACAAGCTGC	GGAGGTTCAA	GCCCAGCTCT	CTGCCGCATC	CGCTCGAGGG
	951	CTCCAGGAAG	GTGGAGGAGG	AAGGAAGCCC	CGGGGACCCC	GACCACGAGG
	1001	CCAGCACCCA	GGGTCGGACC	TGTGAGCCAG	AGCATAGCAA	GGGTGGGGC
	1051	AGGGTGGACG	AGGGGCCCA	GCCACGGAGC	GTGGAGCCCC	AGGATGCGGG
15	1101	ACCCCTGGAG	AGGAGGCAGG	GGGATGAGGC	AGGGGGCCAC	GGGGAAGATA
	1151	GGCCGGAGCC	сттальфссс	AAAGAGAGCA	AGAAGAGGAA	GCTGGAGCTG
	1201	AGCCGGCGGG	AGCAGCCGCC	CACAGGGCCA	GGCCCTCAGA	GTGCCTCAGA
	1251	GGTGGAGAAG	ATCGCTCTGA	ATTTGGAGGG	GTGTGCCCTC	AGCCAGGGCA
	1301	GCCTCAGGAC	GGGGACCCAG	GAAGTGGGCG	GTCAGGACCC	TGGGGAGGCA
20	1351	GTGCAGCCCT	GCCGCCAACC	CCTGGGAGCC	AGGGTGGCCG	ACAAGGTGAG
	1401	GAAGCGGAGG	AAGGTGGATG	AGGGTACTGG	GGACAGTGCT	GCGGTGGCCA
	1451	GTGGTGGTGC	CCAGACCTTG	д ссттессе	GGTCCCCTGC	CCCATCGGGG
	1501	CACCCCAAGG	CTGGACACAG	TGAGAACGGG	GTTGAGGAGG	ACACAGAAGG
<i>'</i>	1551	TCGAACGGGG	CCCAAAGAAG	GTACCCTGG	GAGCCCATCG	GAGACCCCAG
25	1601	GCCCAGCCC	AGCAGGACCT	GCAGGGGACG	AGCCAGCCAA	GACCCCATCG
1	1651	GAGACCCCAG	GCCCAGCCC	GGCAGGACCT	ACAAGGGATG	AGCCAGCCGA
7	1701	GAGCCCATCG	GAGACCCCAG	cccccccccccccccccccccccccccccccccccccc	GGCAGGACCT	GCAGGGGACG
	1751	AGCCAGCCGA	GAGCCCATCG	GAGACCCCAG	GCCCCGCCC	GGCAGGACCT
	1801	GCAGGGGACG	AGCCAGCCAA	GATCCCATC	GAGACCCCAG	GCCCCAGCCC
30	1851	GGCAGGACCT	ACAAGGGATG	AGCCAGCCGA	GAGCCCATCG	GAGACCCCAG
	1901	GCCCCGCCC	GGCAGGACCT	GCAGGGGACG	*GCCAGCCGA	GAGCCCATCG
	1951	GAGACCCCAG	GCCCCGCCC	GGCAGGACCT	GAGGGGACG	AGCCAGCCGA

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		1				
	2001	GAGCCCATCG	GAGACCCCAG	GCCCCAGCCC	GGCAGGACCT	ACAAGGGATG
	2051	AGCCAGCCA	GGCGGGGGAG	GCAGCAGAGT	TGCAGGACGC	AGAGGTGGAG
	2101	TCTTCTGCCA	AGTCTGGGAA	GCCTTAAGGA	AAGGAGTGCC	CGTCGGCGTC
	2151	TTGGTCCTCC	TGTCCCTGCT	GCAGGGGCTG	GGGCCTCCGG	AGCTGCTGCG
5	2201	GGCTCCCCTC	AGGCTCTGCT	TCGTGACCCG	TGACCCATGA	CCCACAGTGC
	2251	TGGCCTCCTG	FGGGGCCACT	ATAGCAGCCA	CCAGAAGCCG	CGAGGCCCTC
	2301	AGGGAAGCCC	AAGGCCTGCA	GAAGCCTCCT	GGCCTGGCTG	TGTCTTCCCC
	2351	ACCCAGCTCT	cdccrececc	CCTGTCTTTG	TAAATTGACC	${\tt CTTCTGGAGT}$
	2401	GGGGGGCG				

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SEQ ID NO: 6 -- Human Opibid Growth Factor Receptor protein, from spliced cDNA version 8

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MDDPDCDSTWEEDEEDAEDAEDEDCEDGEAAGARDADAGDEDEESEEPRAARPSSFOSRM 60 TGSRNWRATRDMCRYRHNYPDLVERDCNGDTPNLSFYRNEIRFLPNGCFIEDILQNWTDN 120 YDLLEDNHSY1QWLFPLREPGVNWHAK#LTLREVEVFKSSQE1QERLVRAYELMLGFYG1 180 RLEDRGTGTVGRAQNYQKRFQNLNWRSHNNLRITRILKSPCELSLEHFQAPLVRFFLEET 240 LVRRELPGVRQSALDYFMFAVRCRHQRRQLVHFAWEHFRPRCKFVWGPQDKLRRFKPSSL 300 PHPLEGSRKVEEEGSPGDPDHEASTQGRTQEPEHSKGGGRVDEGPQPRSVEPQDAGPLER 360 SQGDEAGGHGEDRPEPLSPKESKKRKLELSRREQPPTGPGPQSASEVEKIALNLEGCALS 420 QGSLRTGTQEVGGQDPGEAVQPCRQPLGAR\(\)ADKVRKRRKVDEGTGDSAAVASGGAQTLA 480 LAGSPAPSGHPKAGHSENGVEEDTEGRTGPK GTPGSPSETPGPSPAGPAGDEPAKTPSE 540 TPGPSPAGPTRDEPAESPSETPGPRPAGPAGDEPAESPSETPGPRPAGPAGDEPAKIPSE 600 TPGPSPAGPTRDEPAESPSETPGPRPAGPAGDEPAESPSETPGPRPAGPAGDEPAESPSE 660 697 TPGPSPAGPTRDEPAKAGEAAELQDAEVESSAKGKP

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SEQ ID NO: 7 -- Human OGFr cDNA, spliced version 1

- 1 TAGAATTCAG CGGCCGCTGA ATTCTACCCG AGCATGGACG ACCCCGACTG
- 51 CGACTCCACC TGGGAGGAGG ACGAGGAÇGA TGCGGAGGAC GCGGAGGACG

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SEQ ID NO: 8 -- Human Opioid Growth Factor Receptor protein, from spliced cDNA version 1

- 60 5 MDDPDCDSTWEEDEEDAEDAEDEDCEDGEAAGARDADAGDEDEESEEPRAARPSSFQSRM TGSRNWRATRDMCRYRHNYPDLVERDCNGDTPNLSFYRNEIRFLPNGCFIEDILQNWTDN 120 180 YDLLEDNHSYIQWLFPLREAGVNWHAKPLTLREVEVFKSSQEIQERLVRAYELMLGFYGI RLEDRGTGTVGRAQNYQKRF@NLNWRSHNNLRITRILKSPCELSLEHFQAPLVRFFLEET 240 $\verb|LVRRELPGVRQSALDYFMFAV|| RCRHQRRQLVHFAWEHFRPRCKFVWGPQDKLRRFKPSSL||$ 300 360 10 PHPLEGSRKVEEEGSPGDPDHEASTQGRTCGPEHSKGGGRVDEGPQPRSVEPQDAGPLER SQGDEAGGHGEDRPEPLSPKESKKRKLELSRREQPPTEPGPQSASEVEKIALNLEGCALS 420 461 QGSLRTGTQEVGGQDPGEASCPCqRGWGLRSCCGLPSGSAS
- 15 SEQ ID NO: 9 -- Human OGFT cDNA, spliced version 4
 - 1 TAGAATTCAG CGGCCGCTGA ATTCTAGCCG AGCATGGACG ACCCCGACTG 51 CGACTCCACC TGGGAGGAGG ACGAGGAGGA TGCGGAGGAC GCGGAGGACG 101 AGGACTGCGA GGACGGCGAG GCCGCCGGCG CGAGGGACGC GGACGCAGGG 151 GACGAGGACG AGGAGTCG&A GGAGCCGCGG GCGGCGCGGC CCAGCTCGTT 201 CCAGTCCAGA ATGACAGGGT CCAGAAACTG GCGAGCCACG AGGGACATGT GTAGGTATCG GCACAACTAT CCGGATCTGG TGGAACGAGA CTGCAATGGG 251 301 GACACGCCAA ACCTGAGTTT\CTACAGAAAT GAGATCCGCT TCCTGCCCAA 351 CGGCTGTTTC ATTGAGGACA TTCTTCAGAA CTGGACGGAC AACTATGACC TCCTTGAGGA CAATCACTCC TACATCCAGT GGCTGTTTCC TCTGCGAGAA 401 CCAGGAGTGA ACTGGCATGC CAAGCCCCTC ACGCTCAGGG AGGTCGAGGT 451 501 GTTTAAAAGC TCCCAGGAGA TCQAGGAGCG GCTTGTCCGG GCCTACGAGC TCATGCTGGG CTTCTACGGG ATCCGGCTGG AGGACCGAGG CACGGGCACG 551 GTGGGCCGAG CACAGAACTA CCAGAAGCGC TTCCAGAACC TGAACTGGCG 601 651 CAGCCACAC AACCTCCGCA TCACACGCAT CCTCAAGTCG CCGTGTGAGC TGAGCCTCGA GCACTTCCAG GCGCCACTGG TCCGCTTCTT CCTGGAGGAG 701 751 ACGCTGGTGC GGCGGGAGCT GCCGGGGGGTG CGGCAGAGTG CCCTGGACTA

		1				
	801	CTTCATGTTC	GCCGTGCGCT	GCCGACACCA	GCGCCGCCAG	CTGGTGCACT
	851	TCGCCTGGGA	GCACTTCCGG	CCCCGCTGCA	AGTTCGTCTG	GGGGCCCCAA
	901	GACAAGCTGC	GGAGGTTCAA	GCCCAGCTCT	CTGCCGCATC	CGCTCGAGGG
	951	CTCCAGGAAG	GTGGAGGAGG	AAGGAAGCCC	CGGGGACCCC	GACCACGAGG
5	1001	CCAGCACCCA	GGGTCGGACC	TGTGGGCCAG	AGCATAGCAA	GGGTGGGGC
	1051	AGGGTGGACG	AGGGCCCCA	GCCACGGAGC	GTGGAGCCCC	AGGATGCGGG
	1101	ACCCCTGGAG	AGGAGCCAGG	GGGATGAGGC	AGGGGGCCAC	GGGGAAGATA
	1151	GGCCGGAGCC	dTTAAGCCCC	AAAGAGAGCA	AGAAGAGGAA	GCTGGAGCTG
	1201	AGCCGGCGGG	ACAGCCGCC	CACAGAGCCA	GGCCCTCAGA	GTGCCTCAGA
10	1251	GGTGGAGAAG	ATCGCTCTGA	ATTTGGAGGG	GTGTGCCCTC	AGCCAGGGCA
	1301	GCCTCAGGAC	GGGACCCAG	GAAGTGGGCG	GTCAGGACCC	TGGGGAGGCA
	1351	GTGCAGCCCT	GCCGCCAACC	CCTGGGAGCC	AGGGTGGCCG	ACAAGGTGAG
	1401	GAAGCGGAGG	AAGGTGGATG	AGGGTGCTGG	GGACAGTGCT	GCGGTGGCCA
	1451	GTGGTGGTGC	CCAGACTTG	GCCCTTGCCG	GGTCCCCTGC	CCCATCGGGG
15	1501	CACCCCAAGG	CTGGAQACAG	TGAGAACGGG	GTTGAGGAGG	ACACAGAAGG
	1551	TCGAACGGGG	CCCAAAGAAG	GTACCCCTGG	GAGCCCATCG	GAGACCCCAG
	1601	GCCCCAGCCC	AGCAGGACCT	GCAGGGGACG	AGCCAGCCGA	GAGCCCATCG
	1651	GAGACCCCAG	eccccedcc	AGCAGGACCT	GCAGGGGACG	AGCCAGCCGA
	1701	GAGCCCATCG	GAGACCCCAG	GCCTCCGCCC	GGCAGGACCT	GCAGGGGACG
20	1751	AGCCAGCCGA	GACCCCATCG	GAGACCCCAG	GCCCCAGCCC	GGCAGGACCT
	1801	ACAAGGGATG	AGCCAGCCGA	GAGCCCATCG	GAGACCCCAG	GCCCCGCCC
	1851	GGCAGGACCT	GCAGGGGACG ¹	AGCCAGCCGA	GAGCCCATCG	GAGACCCCAG
	1901	GCCCCGCCC	GGCAGGACCT	CAGGGGACG	AACCAGCCGA	GAGCCCATCG
	1951	GAGACCCCAG	GCCCCAGCCC	GCAGGACCT	ACAAGGGATG	AGCCAGCCAA
25	2001	GGCGGGGGAG	GCAGCAGAGT	TOCAGGACGC	AGAGGTGGAG	TCTTCTGCCA
	2051	AGTCTGGGAA	GCCTTAAGGA	AAGAGTGCC	CGTCGGCGTC	TTGGTCCTCC
X/	2101	TGTCCCTGCT	GCAGGGGCTG	GGGCTCCGG	AGCTGCTGCG	GACTCCCCTC
10/	2151	AGGCTCTGCT	TCGTGACCCG	TGACCCATGA	CCCACAGTGC	TGGCCTCCTG
/	2201	TGGGGCCACT	ATAGCAGCCA	CCAGAAGCCG	CGAGGCCCTC	AGGGAAGCCC
30	2251	AAGGCCTGCA	GAAGCCTCCT	GGCT GGCTG	TGTCTTCCCC	ACCCAGCTCT
	2301	CCCCTGCGCC	CCTGTCTTTG	TAAATTGACC	CTTCTGGAGT	GGGGGGCG
				•		

SEQ ID NO: 10 - Human OGFr, from spliced cDNA version 4

MDDPDCDSTWEEDEEDAEDAEDEDCEDGEAAGARDADAGDEDEESEEPRAARPSSFQSRM	60
TGSRNWRATRDMCRYRHNYPDLVERDCNGDTPNLSFYRNEIRFLPNGCFIEDILQNWTDN	120
YDLLEDNHSYIQWLFP1REPGVNWHAKPLTLREVEVFKSSQEIQERLVRAYELMLGFYGI	180
RLEDRGTGTVGRAQNYQKRFQNLNWRSHNNLRITRILKSPCELSLEHFQAPLVRFFLEET	240
LVRRELPGVRQSALDYFMFAVRCRHQRRQLVHFAWEHFRPRCKFVWGPQDKLRRFKPSSL	300
PHPLEGSRKVEEEGSPGD PHEASTQGRTCGPEHSKGGGRVDEGPQPRSVEPQDAGPLER	360
sqgdeagghgedrpeplspkeskkrklelsrreqpptepgpqsasevekialnlegcals	420
QGSLRTGTQEVGGQDPGEAVQPCRQPLGARVADKVRKRRKVDEGAGDSAAVASGGAQTLA	480
LAGSPAPSGHPKAGHSENGVEEDTEGRTGPKEGTPGSPSETPGPSPAGPAGDEPAESPSE	540
TPGPRPAGPAGDEPAESPSETRGLRPAGPAGDEPAETPSETPGPSPAGPTRDEPAESPSE	600
TPGPRPAGPAGDEPAESPSETPGPRPAGPAGDEPAESPSETPGPSPAGPTRDEPAKAGEA	660
AELQDAEVESSAKSGKP	

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SEQ ID NO: 11 -- Human OGFr cDNA, spliced version 7

TAGAATTCAG CGGCCGCTGA ATTCTAGCCG AGCATGGACG ACCCCGACTG 1 51 CGACTCCACC TGGGAGQAGG ACGAGGAGGA TGCGGAGGAC GCGGAGGACG 101 AGGACTGCGA GGACGGCGAG GCCGCCGGCG CGAGGGACGC GGACGCAGGG 151 201 CCAGTCCAGA ATGACAGGGT CCAGAAACTG GCGAGCCACG AGGGACATGT GTAGGTATCG GCACAACTAT CCGGATCTGG TGGAACGAGA CTGCAATGGG 251 301 GACACGCCAA ACCTGAGTTT\CTACAGAAAT GAGATCCGCT TCCTGCCCAA 351 CGGCTGTTTC ATTGAGGACA TTCTTCAGAA CTGGACGGAC AACTATGACC TCCTTGAGGA CAATCACTCC TACATCCAGT GGCTGTTTCC TCTGCGAGAA 401 451 CCAGGAGTGA ACTGGCATGC CAAGCCCCTC ACGCTCAGGG AGGTCGAGGT 501 GTTTAAAAGC TCCCAGGAGA TCCAGGAGCG GCTTGTCCGG GCCTACGAGC TCATGCTGGG CTTCTACGGG ATCCGGCTGG AGGACCGAGG CACGGGCACG 551 601 GTGGGCCGAG CACAGAACTA CCAGAAGCGC TTCCAGAACC TGAACTGGCG 651 CAGCCACAAC AACCTCCGCA TCACACGCAT CCTCAAGTCG CCGTGTGAGC

			1				
		701	TGAGCCTCGA	GCACTTCCAG	GCGCCACTGG	TCCGCTTCTT	CCTGGAGGAG
		751	ACGCTGGTGC	GGCGGGAGCT	GCCGGGGGTG	CGGCAGAGTG	CCCTGGACTA
		801	CTTCATGTTC	GCCGTGCGCT	GCCGACACCA	GCGCCGCCAG	CTGGTGCACT
		851	TCGCTGGGA	GCACTTCCGG	CCCCGCTGCA	AGTTCGTCTG	GGGCCCCAA
	5	901	GACAAGCTGC	GGAGGTTCAA	GCCCAGCTCT	CTGCCCCATC	CGCTCGAGGG
		951	CTCCAGGAAG	GTGGAGGAGG	AAGGAAGCCC	CGGGGACCCC	GACCACGAGG
		1001	CCAGCACCA	GGGTCGGACC	TGTGGGCCAG	AGCATAGCAA	GGGTGGGGC
		1051	AGGGTGGACG	AGGGGCCCCA	GCCACGGAGC	GTGGAGCCCC	AGGATGCGGG
		1101	ACCCCTGGAG	AGGAGCCAGG	GGGATGAGGC	AGGGGCCAC	GGGGAAGATA
	10	1151	GGCCGGAGCQ	CTTAAGCCCC	AAAGAGAGCA	AGAAGAGGAA	GCTGGAGCTG
		1201	AGCCGGCGGG	AGCAGCCGCC	CACAGAGCCA	GGCCCTCAGA	GTGCCTCAGA
		1251	GGTGGAGAAG	ATCGCTCTGA	ATTTGGAGGG	GTGTGCCCTC	AGCCAGGGCA
		1301	GCCTCAGGAC	GGGACCCAG	GAAGTGGGCG	GTCAGGACCC	TGGGGAGGCA
		1351	GTGCAGCCCT	GCCGCCAACC	CCTGGGAGCC	AGGGTGGCCG	ACAAGGTGAG
	15	1401	GAAGCGGAGG	AAGGTGGATG	AGGGTGCTGG	GGACAGTGCT	GCGGTGGCCA
		1451	GTGGTGGTGC	CCAGACCTTG	GCCCTTGCCG	GGTCCCCTGC	CCCATCGGGG
		1501	CACCCCAAGG	CTGGACACAG	TGAGAACGGG	GTTGAGGAGG	ACACAGAAGG
State of the state	20	1551	TCGAACGGGG	CCCAAAGAAG	GTACCCCTGG	GAGCCCATCG	GAGACCCCAG
- 		1601	GCCCCAGCCC	AGCAGGACCT	GCAGGGGACG	AGCCAGCCGA	GAGCCCATCG
€;		1651	GAGACCCCAG	eccccecc/c	GGCAGGACCT	GCAGGGGACG	AGCCAGCCGA
		1701	GAGCCCATCG	GAGACCCCAG\	GCCCCAGCCC	GGCAGGACCT	ACAAGGGATG
		1751	AGCCAGCCGA	GAGCCCATCG	AGACCCCAG	GCCCCGCCC	GGCAGGACCT
freig freight		1801	GCAGGGGACG	AGCCAGCCGA	GAGCCCATCG	GAGACCCCAG	GCCCCGCCC
		1851	GGCAGGACCT	GCAGGGGACG	AGCAGCCGA	GAGCCCATCG	GAGACCCCAG
	25	1901	GCCCCAGCCC	GGCAGGACCT	ACAAGGGATG	AGCCAGCCAA	GGCGGGGGAG
Dur.		1951	GCAGCAGAGT	TGCAGGACGC	AGAGGTGGAG	TCTTCTGCCA	AGTCTGGGAA
1 (1)		2001	GCCTTAAGGA	AAGGAGTGCC	CGTCGGCGTC	TTGGTCCTCC	TGTCCCTGCT
		2051	GCAGGGGCTG	GGGCCTCCGG	AGCTGCT CG	GGCTCCCCTC	AGGCTCTGCT
0/		2101	TCGTGACCCG	TGACCCATGA	CCCACAGTC	TGGCCTCCTG	TGGGGCCACT
	30	2151	ATAGCAGCCA	CCAGAAGCCG	CGAGGCCCT	AGGGAAGCCC	AAGGCCTGCA
•		2201	GAAGCCTCCT	GGCCTGGCTG	TGTCTTCCCC \	ACCCAGCTCT	CCCCTGCGCC
		2251	CCTGTCTTTG	TAAATTGACC	CTTCTGGAGT	deeeecee	



SEQ ID NO: 12, Human OGFr, from spliced cDNA version 7

MDDPDCDSTWEEDEEDAEDAEDEDCEDGEAAGARDADAGDEDEESEEPRAARPSSFQSRM	60
TGSRNWRATRDMCRYRHNYPDLVERDCNGDTPNLSFYRNEIRFLPNGCFIEDILQNWTDN	120
YDLLEDNHSYIQWLFPLREPGVNWHAKPLTLREVEVFKSSQEIQERLVRAYELMLGFYGI	180
RLEDRGTGTVGRAQNYOKRFQNLNWRSHNNLRITRILKSPCELSLEHFQAPLVRFFLEET	240
LVRRELPGVRQSALDYFMFAVRCRHQRRQLVHFAWEHFRPRCKFVWGPQDKLRRFKPSSL	300
PHPLEGSRKVEEEGSPG PDHEASTQGRTCGPEHSKGGGRVDEGPQPRSVEPQDAGPLER	360
SQGDEAGGHGEDRPEPLS KESKKRKLELSRREQPPTEPGPQSASEVEKIALNLEGCALS	420
QGSLRTGTQEVGGQDPGEAVQPCRQPLGARVADKVRKRRKVDEGAGDSAAVASGGAQTLA	480
LAGSPAPSGHPKAGHSENGVEEDTEGRTGPKEGTPGSPSETPGPSPAGPAGDEPAESPSE	540
TPGPRPAGPAGDEPAESPSETPGPSPAGPTRDEPAESPSETPGPRPAGPAGDEPAESPSE	600
TPGPRPAGPAGDEPAESPSET PGPSPAGPTRDEPAKAGEAAELQDAEVESSAKSGKP	657

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SEQ ID NO: 13 -- Human OGFr cDNA, spliced version 127

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51 CGACTCCACC TGGGAGAGG ACGAGGAGGA TGCGGAGGAC GCGGAGGACG
101 AGGACTGCGA GGACGGGGAG GCCGCCGGCG CGAGGGACGC GGACGCAGGG

151 GACGAGGACG AGGAGTCGGA GGAGCCGCGG GCGGCGCGCC CCAGCTCGTT

1 TAGAATTCAG CGGCCGCTGA ATTCTAGCCG AGCATGGACG ACCCCGACTG

201 CCAGTCCAGA ATGACAGGT CCAGAAACTG GCGAGCCACG AGGGACATGT

251 GTAGGTATCG GCACAACTAT CCGGATCTGG TGGAACGAGA CTGCAATGGG

301 GACACGCCAA ACCTGAGTTT\CTACAGAAAT GAGATCCGCT TCCTGCCCAA
351 CGGCTGTTTC ATTGAGGACA\TTCTTCAGAA CTGGACGGAC AACTATGACC

401 TCCTTGAGGA CAATCACTCC TACATCCAGT GGCTGTTTCC TCTGCGAGAA

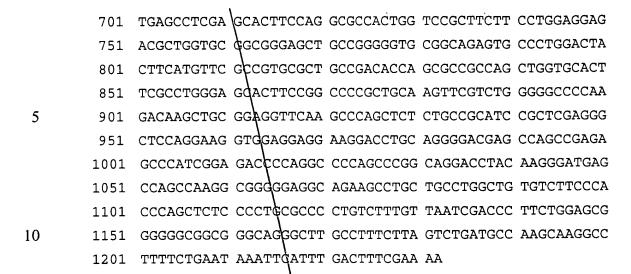
451 CCAGGAGTGA ACTGGCATGC QAAGCCCCTC ACGCTCAGGG AGGTCGAGGT

501 GTTTAAAAGC TCCCAGGAGA TQCAGGAGCG GCTTGTCCGG GCCTACGAGC

551 TCATGCTGGG CTTCTACGGG ATCCGGCTGG AGGACCGAGG CACGGGCACG

601 GTGGGCCGAG CACAGAACTA CCAGAAGCGC TTCCAGAACC TGAACTGGCG

651 CAGCCACAAC AACCTCCGCA TCACACGCAT CCTCAAGTCG CCGTGTGAGC



SEQ ID NO: 14 -- Human OGFr, from spliced cDNA version 127

MDDPDCDSTWEEDEEDAEDAEDAEDCEDGEAAGARDADAGDEDEESEEPRAARPSSFQSRM

TGSRNWRATRDMCRYRHNYPDLVERDCNGDTPNLSFYRNEIRFLPNGCFIEDILQNWTDN

120

YDLLEDNHSYIQWLFPLREPGVNWHAKPLTLREVEVFKSSQEIQERLVRAYELMLGFYGI

RLEDRGTGTVGRAQNYQKRFQNLNWRSHNNLRITRILKSPCELSLEHFQAPLVRFFLEET

240

LVRRELPGVRQSALDYFMFAVRCRHQRRQLVHFAWEHFRPRCKFVWGPQDKLRRFKPSSL

300

PHPLEGSRKVEEEGPAGDEPAESPSETPGPSPAGPTRDEPAKAGEAEACCLAVSSHPALP

360

CAPVFVNRPFWSGGRRAGLAFLSLMPSKAFSE